

# SEQUENCE LISTING

SEQ ID NO: 1:

GGCTCCTCATCTGGAACACCTCGGGTCACCCCCGACAACGGTGGTGGGAGGGAGAGCGGC	60
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TGGGTTTGCCCCAATGACCGGCAGCTTGCCCTTCGAGCCAAGCTGCAGACGGGCTGGTCC	240
GTGCACACCTACCAGACGGAGAAGCAGAGGAGGAAGCAGCACCTCAGCCCGCGGAGGTG	300
GAGGCCATCCTGCAGGTCATCCAGAGGGCAGAGCGGCTCGACGTCCTGGAGCAGCAGAGA	360
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TCCCAGTGTCTGCTCTGCGGGGAGGTGCTGGGCTTCCTGGGCAGCTCGTCGGTGTCTGC	480
AAAGACTGCAGGAAGGTCTGGAAGAGGTGGGGGGCCTGGTTCTACAAAGGGCTCCCCAAG	540
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GAGGACAGACTCCCATCCACTGGGGTCAGGGACCGGAAGGCGACAAACCCTGGAAGGAG	780
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CTCCGAAGCTGCGTGTGGCCCCCTTAGAGGTGAGCATCAGAGCCAGAGCAGTGAGGGGGAG	1500
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TGTTTGTGTTGTTTTGACACAGTCTCGCTTTGTTGCCCAGGCTGGGGTGCAGTGGCACGA	1680
TCGGGCTCACTGCAACCTCCACCTCCCGGGCTCAAGCGATTCTCTCACCTCAGCCTCCT	1740
GAGTAGGTGGGATTACAGATGCCCGCCACCACCCAGTTAATTTTGTATTTTGAAG	1800
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SEQ ID NO: 2:

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 ArgAlaGluArgLeuAspValLeuGluGlnGlnArgIleGlyArgLeuValGluArgLeu 80  
 GluThrMetArgArgAsnValMetGlyAsnGlyLeuSerGlnCysLeuLeuCysGlyGlu 100  
 ValLeuGlyPheLeuGlySerSerSerValPheCysLysAspCysArgLysValTrpLys 120  
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 GlyArgAlaAspGluProGlnPheArgProTrpProThrGluProAlaGluArgGluPro 160  
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 SerAspSerAspSerAspLeuSerSerSerSerLeuGluAspArgLeuProSerThrGly 200  
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 ArgMetGlyPheThrGlnProAlaGlyHisLeuPheGlyLeuGlnSerSerLeuAlaSer 240  
 GlyGluThrGlyThrGlySerAlaAspProProGlyGlyGlyThrGlySerAlaAspPro 260  
 ProGlyGlyProArgProGlyLeuThrArgArgAlaProValLysAspThrProGlyArg 280  
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SEQ ID NO: 3:

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 GAGGCCATCCTGCAGGTCATCCAGAGGGCAGAGCGGCTCGACGTCTGGAGCAGCAGAGA 360  
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CATCAGAGCCAGAGCAGTGAGGGGGAGACTCACCCACCTCTCCCTCTCCCTTCAGCTCT	1620
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CACTTGCTGAAGAGCAGCGTTTCAAGTGCATCCCCAGCCAGGGCACGTGGCTCCCTCAGCC	2160
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SEQ ID NO: 4:

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ArgAlaGluArgLeuAspValLeuGluGlnGlnArgIleGlyArgLeuValGluArgLeu 80  
GluThrMetArgArgAsnValMetGlyAsnGlyLeuSerGlnCysLeuLeuCysGlyGlu 100  
ValLeuGlyPheLeuGlySerSerSerValPheCysLysAspCysArgLysLysValCys 120  
ThrLysCysGlyIleGluAlaSerProGlyGlnLysArgProLeuTrpLeuCysLysIle 140  
CysSerGluGlnArgGluValTrpLysArgSerGlyAlaTrpPheTyrLysGlyLeuPro 160  
LysTyrIleLeuProLeuLysThrProGlyArgAlaAspAspProHisPheArgProLeu 180  
ProThrGluProAlaGluArgGluProArgSerSerGluThrSerArgIleTyrThrTrp 200  
AlaArgGlyArgValValSerSerAspSerAspSerAspSerAspLeuSerSerSerSer 220  
LeuGluAspArgLeuProSerThrGlyValArgAspArgLysGlyAspLysProTrpLys 240  
GluSerGlyGlySerValGluAlaProArgMetGlyPheThrGlnProAlaGlyHisLeu 260  
PheGlyLeuGlnSerSerLeuAlaSerGlyGluThrGlyThrGlySerAlaAspProPro 280  
GlyGlyGlyThrGlySerAlaAspProProGlyGlyProArgProGlyLeuThrArgArg 300  
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SerSerCysLeuGly 325

SEQ ID NO: 5:

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GACAGTGAAGTCCGATCTTAGCTCCTCCAGCCTAGAGGACAGACTCCCATCCACTGGGGTC 900

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TGAACCTCACTTGCTGAAGAGCAGCGTTCAGGTGCATCCCCAGCCAGGGCACGTGGCTCCC	2220
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AGTATGGGCAAATGCTTCTGAAAACCCCTTCCCTGAAGAGAGAGAACGTGTGTGTGTGTG	2340
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SEQ ID NO: 6:

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CysGlyIleGluAlaSerProGlyGlnLysArgProLeuTrpLeuCysLysIleCysSer	60
GluGlnArgGluValTrpLysArgSerGlyAlaTrpPheTyrLysGlyLeuProLysTyr	80
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SEQ ID NO: 7:

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TGAAGATTAAAA	2592

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